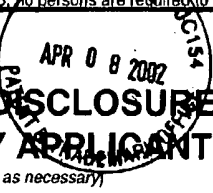


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			Application Number			
			Filing Date		August 20, 2001	
			First Named Inventor		Person, Christophe	
			Group Art Unit		Not Yet Assigned 1631	
Examiner Name		Not Yet Assigned J. Brosca				
Sheet	1	of	1	Attorney Docket Number		LXGN-00104

U.S. PATENT DOCUMENTS						
Examiner Initials*	Cite No. ¹	U.S. Patent Document		Name of Patentee or Applicant of Cited Document	Date of Publication of cited Document MM-DD-YYYY	Pages, Columns, Lines, Where Relevant Passages or Relevant Figures Appear
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JLB	AA	ALTSCHUL, STEPHEN F. ET AL, 1990, "Basic Local Alignment Search Tool", J. Mol. Biol. 215:403-410.			
JLB	AB	ALTSCHUL, STEPHEN F. ET AL, 1997, "Gapped BLAST and PSI-BLAST: a new generation of protein database search programs", Nucleic Acids Res. 25:3389-3402.			
JLB	AC	FENG, F. ET AL, 1984-85, "Aligning amino acid sequences: comparison of commonly used methods", J Mol Evol. 21(2): 112-25.			
JLB	AD	HENIKOFF S., and HENIKOFF, J.G., 1992, "Amino acid substitution matrices from protein blocks", Proc Natl Acad Sci USA 89(22):10915-9.			
JLB	AE	KARLIN, S. and GHANDOUR, G., 1985, "Multiple-alphabet amino acid sequence comparisons of the immunoglobulin kappa-chain constant domain", Proc Natl Acad Sci USA 82(24):8597-601.			
JLB	AF	LIPMAN, DAVID J. and PEARSON, W.R., 1985, "Rapid and sensitive similarity searches", Science 227:1435-1441.			
JLB	AG	PEARSON, W. and LIPMAN, DAVID, 1998, "Improved tools for biological sequence comparison", Proc. Natl. Acad. Sci. 85:2444-2448.			
JLB	AH	PEARSON, W., 1990, "Rapid and sensitive sequence comparison with FASTP and FASTA", Methods in Enzymology 183:63-98.			
JLB	AI	SMITH, T.F. and WATERMAN, M.S., 1981, "Identification of common molecular subsequences", J. Mol. Biol. 147:195-197.			

Examiner Signature	JLB. Brosca	Date Considered	20 July 2004
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